

## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/996,128

Source: OIPE

Date Processed by STIC: 12/6/01

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**

**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**

- 1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

**FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.**

**FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.**

**PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)**

**PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)**

**TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:**

### **Checker Version 3.0**

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

**Checker Version 3.0 can be down loaded from the USPTO website at the following address:**

**<http://www.uspto.gov/web/offices/pac/checker>**

## Raw Sequence Listing Error Summary

ERROR DETECTED    SUGGESTED CORRECTION    SERIAL NUMBER: 09/996,128

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1  Wrapped Nucleic  
     Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2  Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.

3  Misaligned Amino  
    Numbering The numbering under each 5<sup>th</sup> amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

4  Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

5  Variable Length Sequence(s) \_\_\_\_\_ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6  PatentIn 2.0  
    "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) \_\_\_\_\_. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

7  Skipped Sequences  
    (OLD RULES) Sequence(s) \_\_\_\_\_ missing. If intentional, please insert the following lines for each skipped sequence:  
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
(i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)  
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

8  Skipped Sequences  
    (NEW RULES) Sequence(s) \_\_\_\_\_ missing. If intentional, please insert the following lines for each skipped sequence.  
<210> sequence id number  
<400> sequence id number  
000

9  Use of n's or Xaa's  
    (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.  
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.  
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10  Invalid <213>  
    Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

11  Use of <220> Sequence(s) \_\_\_\_\_ missing the <220> "Feature" and associated numeric identifiers and responses.  
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.  
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

12  PatentIn 2.0  
    "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

OIPE

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/996,128

DATE: 12/06/2001

TIME: 11:42:50

Input Set : A:\sequence listing.txt  
 Output Set: N:\CRF3\12062001\I996128.raw

Does Not Comply  
 Corrected Diskette Needed

Errors on pp. 1,3

3 <110> APPLICANT: Houghton, Alan  
 4 Bergman, Phillip  
 5 Wolchok, Jedd  
 7 <120> TITLE OF INVENTION: Compositions for treatment of Melanoma and Methods of Using  
 Same  
 9 <130> FILE REFERENCE: MSK.P-026-3  
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/996,128  
 C--> 11 <141> CURRENT FILING DATE: 2001-11-27  
 11 <150> PRIOR APPLICATION NUMBER: US 09/627,694  
 12 <151> PRIOR FILING DATE: 2000-07-28  
 14 <150> PRIOR APPLICATION NUMBER: US 09/308,697  
 15 <151> PRIOR FILING DATE: 1999-05-21  
 17 <150> PRIOR APPLICATION NUMBER: PCT/US97/22669  
 18 <151> PRIOR FILING DATE: 1997-12-10  
 20 <150> PRIOR APPLICATION NUMBER: US 60/036,419  
 21 <151> PRIOR FILING DATE: 1997-02-18  
 23 <150> PRIOR APPLICATION NUMBER: US 60/032,535  
 24 <151> PRIOR FILING DATE: 1996-12-10  
 26 <150> PRIOR APPLICATION NUMBER: US 60/180,651  
 27 <151> PRIOR FILING DATE: 2000-01-26  
 29 <160> NUMBER OF SEQ ID NOS: 2  
 31 <170> SOFTWARE: PatentIn version 3.0  
 33 <210> SEQ ID NO: 1  
 34 <211> LENGTH: 6408  
 35 <212> TYPE: DNA  
 36 <213> ORGANISM: synthetic construct  
 38 <400> SEQUENCE: 1  
 39 attctgcaga tatccagcac agtggcgcc gctcgagtct agagggcccg tttaaacccg 60  
 41 ctgatcagcc tcgactgtgc ctcttagttt ccagccatct gttgtttgcc cctccccgt 120  
 43 gccttccttg acccttggaaag gtgccactcc cactgtcctt tcctaataaa atgagggaaat 180  
 45 tgcacatcgat tgcgttgc ggtgttgcattc tattctgggg ggtgggggtgg ggcaggacag 240  
 47 caaggggggag gattggaaag acaatagcag gcatgggggg gatgcagggg gggggggggcg 300  
 49 ctgagggtctg cctcgtaag aagggtttgc tgactcatac caggcctgaa tcgccccatc 360  
 51 atccagccag aaagtgggg agccacgggt gatgagagct ttgttgcattt tggaccagtt 420  
 53 ggtgattttt aacttttgc ttgcacggg acggctgcg ttgtcgggaa gatgcgtgat 480  
 55 ctgatccttc aactcagcaa aagtgcatt tattcaacaa agccgcgcgc ccgtcaagtc 540  
 57 agcgtaatgc tctgccatgt ttacaaccaa ttaaccaatt ctgatttagaa aaactcatcg 600  
 59 agcatcaaat gaaactgcaa ttatttcata tcaggattat caataccata tttttgaaaa 660  
 61 agccgtttct gtaatgaagg agaaaaactca ccgaggcagt tccataggat ggcaagatcc 720  
 63 tggatcggt ctgcattcc gactcgatcca acatcaatac aacctattaa ttcccctcg 780  
 65 tcaaaaataa ggttatacaag tgagaaatca ccatgagtga cgactgaatc cggtgagaat 840  
 67 ggcaaaaagct tatgcatttc ttccagact tggtaacaa ggcaggcatt acgctcgatca 900  
 69 tcaaaaatcac tcgcatcaac caaaccgtta ttcattcgat attgcgcctg agcgagacga 960  
 71 aatacgcgtat cgctgttaaa aggacaatta caaacaggaa tcgaatgcaa ccggcgccagg 1020  
 73 aacactgcca ggcgcataac aatattttca cctgaatcag gatatttttc taatacctgg 1080  
 75 aatgcgtttt tcccgggat cgcaatggat agtaaccatg catcatcagg agtacggata 1140  
 77 aatgcgttga tggtcgaaag agcataaaat tccgtcagcc agttagtct gaccatctca 1200  
 79 tctgttaacat cattggcaac gctaccttg ccatgttca gaaacaactc tggcgcatcg 1260

invalid response, see Error summary sheet  
 item 10

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Input Set : A:\sequence listing.txt

Output Set: N:\CRF3\12062001\I996128.raw

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85	tcccgaa tatggctcat aacacccctt gtattactgt ttatgtaaac agacagtttt	1440
87	atgttcatg atgatatatt ttatcttgc gcaatgtaaac atcagagatt ttgagacaca	1500
89	acgtggctt ccccccccccc cctgcagcg ttcttcctt tccccacccc accccccaag	1560
91	ttcgggtgaa gccccaggc tocgagccaa cgtcggggcg gcaggccctg ccatacgctc	1620
93	agttactca tatatactt agattgattt aaaacttcat tttaattta aaaggatcta	1680
95	ggtgaagatc cttttgata atctcatgac caaaatccct taacgtgagt ttctgtcca	1740
97	ctgagcgtca gaccccgtag aaaagatcaa aggatcttgc tagatccctt ttttctgcg	1800
99	cgtaatctgc tgctgcaaa caaaaaaacc accgctacca gcggtgggtt gtttgcgg	1860
101	tcaagagcta ccaactctt ttccgaaggt aactggcttc agcagagcgc agataccaaa	1920
103	tactgttctt ctatgttagc ctagttagg ccaccatc aagaactctg tagcaccgccc	1980
105	tacataccctc gctctgctaa ttctgttacc agtggctgtc gccagtggcg ataagtctg	2040
107	tcttaccggg ttggactcaa gacgatagtt accggataag ggcgcggcgt cggctgaac	2100
109	gggggggttcg tgcacacagc ccagcttggc gcaacgacc tacaccgaaac tgagataacct	2160
111	acagcgttag ctatgagaaa ggcacacgct tcccaagg agaaaggcgg acaggtatcc	2220
113	ggtaagcggc agggctggaa caggagagcg cacgagggag ctccagggg gaaacgcctg	2280
115	gtatctttat agtcctgtcg gtttcgcctt cctctgactt gagcgtcgat ttttgtatg	2340
117	ctcgtcaggg gggcggagcc tatggaaaaa cgccagcaac gccccttt tacggttcct	2400
119	ggcctttgc tggcttttg ctcacatgtt ctccatgtc ttatccctg attctgtgaa	2460
121	taaccgtatt accgcattgc attagttatt aatagtaatc aattacgggg tcattagttc	2520
123	atagccata tatggagttc cgcgttacat aactacggt aatggcccg cttggctgac	2580
125	cggccaaacga ccccccggca ttgacgtcaa taatgacgag atctgatata ggtgacagac	2640
127	gatatgaggc tatatcgccg atagaggcga catcaagctg gcacatggcc aatgcataatc	2700
129	gatctataca ttgaatcaat attggcaatt agccatatta gtcattgggtt atatagcata	2760
131	aatcaatatt ggctattggc cattgcatac gtttatcta tattataata tgtacattta	2820
133	tattggctca tgtccaaat gacccatgt ttgacattga ttattgacta gttattaaata	2880
135	gtaatcaatt acggggctat tagttcatag cccatatatg gagttcccg ttacataact	2940
137	tacggtaat gggccgcctg gctgaccggc caacgacccc cggccatgtg cgtcaatgtat	3000
139	gacgtatgtt cccatagtaa cgcacatagg gacttccat tgacgtcaat gggggagta	3060
141	tttacggtaa actgcccact tggcgttaca tcaagtgtat catatgcacaa gtccggccccc	3120
143	tattgacgtc aatgacggta aatggccgc ctggcattat gcccagtaca tgaccttacg	3180
145	ggactttccct acttggcagt acatctacgt attagtcac gctattacca tgggtatgcg	3240
147	gttttggcag tacaccaatg ggcgtggata gcggttgac tcacggggat ttccaagtct	3300
149	ccacccattt gacgtcaatg ggagttgtt ttggcaccaa aatcaacggg acttccaaa	3360
151	atgtcgtaat aacccggccc ctggacgca aatggccgtt aggcgtgtac gttggggaggt	3420
153	ctatataaggc agagctcgat tagtgaaccc tcagatcgcc tggagacgcc atccacgctg	3480
155	ttttgacctc catagaagac accgggaccc atccacgcctc cgcggccggg aacgggtcat	3540
157	tggacgcgg attcccggtt ccaagatgtg cgttaagtacc gcctatagac tctataggca	3600
159	cacccctttt gctcttatgc atgtataact gttttggct tggggccat acaccccccgc	3660
161	ttccttatgc tataggtat ggtatagctt agcctatagg tttttttat tgaccattat	3720
163	tgaccactcc cctattggtg acgatactt ccattactaa tccataacat ggctctttgc	3780
165	cacaactatc tctattggct atatgcacat actctgtccct tcagagactg acacggactc	3840
167	tgtatttta caggatgggg tcccatatatt tattacaaa ttccatata caacaacgc	3900
169	gtccccctgtt cccgcgttt ttatcaaaca tagcgtggta tctccacgcg aatctgggt	3960
171	acgtgtccg gacatgggtt cttctccgtt agcggcgag cttccacatc cgagccctgg	4020
173	tcccatgcct ccagcggctc atggctgcctc ggcagctctt tgctctaaac agtggaggcc	4080
175	agacttaggc acagcacaat gcccaccacc accagtgtgc cgcacaaggc cgtggcggt	4140
177	gggtatgtt ctgaaaatga gtcggagat tggctcgca ccgctgacgc agatgaaaga	4200

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PATENT APPLICATION: US/09/996,128

DATE: 12/06/2001

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Input Set : A:\sequence listing.txt

Output Set: N:\CRF3\12062001\I996128.raw

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183	gttgctgccg	cgcgcgccac	cagacataat	agctgacaga	ctaacagact	gttccttcc	4380
185	atgggtcttt	tctgcagtca	ccgtccacgc	gtaatacga	ctcaactatag	ggagacccaa	4440
187	gctggctagc	gtttaaactt	aagcttggta	ccgagctcgg	atccactagt	ccagtgtgg	4500
189	ggaattccgg	gaagaatgct	cctggctgtt	ttgtactgcc	tgctgtggag	tttccagacc	4560
191	tccgctggcc	atttccttag	agcctgtgtc	tcctctaaga	acctgatgga	gaaggaatgc	4620
193	tgtccaccgt	ggagcgggg	caggagtccc	tgtggccagc	tttcaggcag	aggttcctgt	4680
195	cagaatatcc	ttctgtccaa	tgcaccactt	gggcctcaat	ttcccttcac	aggggtggat	4740
197	gaccgggagt	cgtggccttc	cgtcttttat	aataggacat	gccagtgtc	tggcaacttc	4800
199	atgggattca	actgtggaaa	ctgcaagttt	ggcttttggg	gaccaaactg	cacagagaga	4860
201	cgactcttgg	tgagaagaaa	catttcgtat	ttgagtgtccc	cagagaagga	caaattttt	4920
203	gcctacactca	ctttagcaaa	gcataaccatc	agctcagact	atgtcatccc	catagggacc	4980
205	tatggccaaa	tgaaaaatgg	atcaacacacc	atgtttaacg	acatcaatat	ttatgacctc	5040
207	tttgcatttgc	tgcatttata	tgtgtcaatg	gatgcactgc	ttgggggata	tgaatctgg	5100
209	agagacattt	attttgcaca	tgaagcacca	gttttctgc	cttggcatag	actcttcttgc	5160
211	ttgcgggtgg	aacaagaaat	ccagaagctg	acagagatg	aaaacttcac	tattccatat	5220
213	tgggactggc	gggatgcaga	aaagtgtgac	atttgcacag	atgagtatcat	gggaggtcag	5280
215	caccccacaa	atcctaactt	actcagccca	gcatcattct	tcccttcgt	gcagattgtc	5340
217	tgtagccgt	tggaggagta	caacagccat	cagtctttat	gcaatggaa	gcccgggg	5400
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223	gccaatttca	gcttagaaa	tacactggaa	ggatttgcata	gtccacttac	tggatagcg	5580
225	gatgcctctc	aaagcagcat	gcacaatgcc	ttgcacatct	atatgaatgg	aacaatgtcc	5640
227	caggtacagg	gatctgccaa	cgatcctata	tcccttcata	accatgcatt	tgttgacagt	5700
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249	tttaaacatt	ttcccttaag	cccatatgtc	taaggaaagg	atgttatttgc	taatgagga	6360
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254	<210>	SEQ ID NO:	2				
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256	<212>	TYPE:	DNA				
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259	<400>	SEQUENCE:	2				
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262	ctgatcagcc	tcgactgtgc	cttctagttt	ccagccatct	gttggggcc	cctccccccgt	120
264	gccttccttgc	acccttggaa	gtgccactcc	cactgtcctt	tcctataaaa	atgagaaat	180
266	tgcatcgcat	tgtctgatgt	ggtgtcattt	tattctgggg	ggtgggggtgg	ggcaggacag	240
268	caagggggag	gattggaaag	acaatagcag	gcatgggg	gatgcaggggg	ggggggggcg	300
270	ctgagggtctg	cctcgtaag	aagggtttgc	tgactcatac	caggcctgaa	tcgccccatc	360
272	atccagccag	aaagtgggg	agccacgg	gatgagagct	ttgtttagg	tggaccagtt	420

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Input Set : A:\sequence listing.txt

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278	agcgtaatgc tctgccagtgt ttacaaccaa ttaaccaatt	ctgatttagaa aaactcatcg	600
280	agcatcaaataat gaaactgcaa tttattcata tcaggattat	caataccata ttttggaaaa	660
282	agccgttct gtaatgaagg agaaaactca ccgaggcagt	tccataggat ggcaagatcc	720
284	tgttatcggt ctgcgattcc gactcgtcca acatcaatac	aacctattaa tttccctcg	780
286	tcaaaaataaa ggttatcaag tgagaaatca ccatgagtga	cgactgaatc cggtgagaat	840
288	gcaaaaagct tatgcatttc tttccagact tgttcaacag	gccagccatt acgctcgta	900
290	tcaaaaatcac tcgcacatcaac caaaccgtta ttcattcg	tttgcgcctg akgagacga	960
292	aatacgcgt cgctgttaaa aggacaatta caaacaggaa	tcgaatgca cccgcgcagg	1020
294	aacactgcca gcgcacatcaac aatatttca cctgaatcg	gatattcttc taatacctgg	1080
296	aatgctgttt tcccgggat cgcaagtggt agtaaccatg	catcatcagg agtacggata	1140
298	aaatgcttga tggtcggaag aggataaat tccgtcagcc	agtttagtct gaccatctca	1200
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304	ttataccatataatcagc atccatgtt gatttaatc	gcggccctcg gcaagacgtt	1380
306	tcccggtt gaa tatggctcat aacaccctt gtattactgt	ttatgtaaagc agacagtttt	1440
308	attgttcatg atgatataattttatcttgc acaatgttgc	atcagagatt ttgagacaca	1500
310	acgtggctt ccccccccccc cctgcagcgt ttcttcctt	tcccccacccc accccccaag	1560
312	tccgggtgaa ggcccaggc tcgcagccaa cgtcgccgc	gcaggccctg ccatagccctc	1620
314	aggtaactca tatatacttt agattgattt aaaacttcat	tttaattta aaaggatcta	1680
316	ggtaagatc cttttgtata atctcatgac caaaatccct	taacgtgagt ttgcgttcca	1740
318	ctgagcgtca gaccccgtag aaaagatcaa aggatcttgc	tgagatcctt ttttctgcg	1800
320	cgtaatctgc tgcttgcaaa caaaaaacc accgtatcca	gcgggtgggtt gttgccgga	1860
322	tcaagagctt ccaactctt ttccgaaggt aactggcttc	agcagagcgc agataccaaa	1920
324	tactgttctt ctatgttgc ctatgttgc ccaccacttc	aagaactctg tagcaccgc	1980
326	tacatacctc gctctgtctaa tcctgttacc agtgcgtct	gccagtgccg ataagtcgt	2040
328	tcttaccggg ttggactcaa gacgatagg accggataag	gcgcagccgt cggctgaac	2100
330	gggggggttcg tgcacacagc ccagcttggaa	gcgaacgacc tacaccgaac tgagatacc	2160
332	acagcgttagatgatgaa ggcgcacgtt tcccaaggaa	agaaaggccg acaggtatcc	2220
334	ggtaagcggc agggtcggaa caggagagcg cacgagggag	cttccaggg gaaacgcctg	2280
336	gtatctttat agtccgtcg gtttcgcctt cctctgactt	gagcgtcgat ttttgcgt	2340
338	ctcgtcaggg gggcgagcc tatggaaaaaa cgcacatcg	gcggcccttt tacgttcc	2400
340	ggcctttgc tggccttttgc ctcacatgtt ctttgcgt	ttatccctg attctgttgc	2460
342	taaccgtatt accgcatttc attagttt aatagtaatc	aattacgggg tcattatgtt	2520
344	atagccata tatggatgtc cgcgttacat aacttacgtt	aatggcccg cctggctgac	2580
346	cgcacatcg cccccccca ttgacgtcaa taatgacgat	atctgtatata ggtgacagac	2640
348	gatatgaggc tatatcgccg atagaggcga catcaagctg	gcacatggcc aatgcataatc	2700
350	gatctataca ttgatcaat attggcaatt agccatatta	gtcattgggtt atatagcata	2760
352	aatcaatatt ggctattggc cattgcatac gttgtatcta	tatcataata tgtacattt	2820
354	tattggctca tgtccatat gaccgcatttgc tgcattgtt	ttattgtactt gttatataat	2880
356	gtaatcaatt acgggtcat tagttcatag cccatataatg	gagttccgcg ttacataact	2940
358	tacggtaat ggcggccctg gtcgaccgc caacgacccc	cgcccatatgtatgcgtat	3000
360	gacgtatgtt cccatagtaa cgcacatagg gacttccat	tgacgtcaat gggggagta	3060
362	tttacggtaa actgcccact tggcgttaca tcaatgtat	catatgcacat gtcggccccc	3120
364	tattgacgtc aatgacggta aatggccgc ctggcattat	gcccagtaca tgaccttacg	3180
366	ggacttcctt acttggcagt acatctacgtt attagtcac	gctattacca tgggtatgcg	3240
368	gttttggcag tacaccaatg ggcgtggata ggcgttgc	tcacggggat ttccaagtct	3300
370	ccacccattt gacgtcaatg ggagtttttttgc tggcaccat	aatcaacggg acatccaaa	3360

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/996,128

DATE: 12/06/2001  
TIME: 11:42:50

Input Set : A:\sequence listing.txt  
Output Set: N:\CRF3\12062001\I996128.raw

372	atgtcgtaat	aaccccgccc	cggtgacgca	aatgggcgg	aggcgtgtac	ggtgggaggt	3420
374	ctatataagc	agagctcggt	tagtgaaccg	tcagatcgcc	tggagacgcc	atccacgctg	3480
376	ttttgacctc	catagaagac	accgggaccc	atccagccctc	cgcggccggg	aacgggtgc	3540
378	tggaacgcgg	atccccgtg	ccaagagtga	cgtaagtacc	gcctataagac	tctataggca	3600
380	cacccctttg	gctcttatgc	atgctatact	gttttggct	tggggcctat	acaccccccgc	3660
382	ttccttatgc	tataggtgat	ggtatagctt	agcctatagg	tgtgggttat	tgaccattat	3720
384	tgaccactcc	cctattgggt	acgatacttt	ccattactaa	tccataacat	ggctctttgc	3780
386	cacaactatc	tctattggct	atatgccaat	actctgtcct	tcagagactg	acacggactc	3840
388	tgtatTTTA	caggatgggg	tcccatttat	tatttacaaa	ttcacatata	caacaacgcc	3900
390	gtccccccgtg	cccgcagttt	ttattaaaca	tagcgtggga	tctccacgcg	aatctcggt	3960
392	acgtgttccg	gacatgggt	cttctccgg	agcggcggag	cttccacatc	cgagccctgg	4020
394	tcccatgcct	ccagcggctc	atggtcgctc	ggcagctcct	tgctcttaac	agtggaggcc	4080
396	agacttaggc	acagcacaat	gccaccacc	accagtgtgc	cgcacaaggc	cgtggcggta	4140
398	gggtatgtgt	ctgaaaatga	gctcgagat	tgggtcgc	ccgctgacgc	agatggaaga	4200
400	cttaaggcag	cggcagaaga	agatgcagc	agctgagttt	ttgtattctt	ataagagtca	4260
402	gaggtaactc	ccgttgcgg	gctgttaacg	gtggagggca	gtgtagtc	agcagtactc	4320
404	gttgctgccc	cgcgcgccac	cagacataat	agctgacaga	ctaacagact	gttccctttcc	4380
406	atgggtcttt	tctgcagtca	cogtccacgc	gttaatacga	ctcaactatag	ggagacccaa	4440
408	gtggctagc	gtttaaactt	aagcttggta	ccgagctcgg	atccactatgt	ccagtgtgg	4500
410	ggaattctgt	gacactcatt	aacattttgg	tgcagatttt	gtatgatcta	aaggagaaaa	4560
412	tgttcttggc	tgtttgtat	tgccttctgt	ggagttcca	gatctctgt	ggcattttc	4620
414	ctcgagccctg	tgcctcctct	aagaacttgg	tggcaaaaga	atgctgccc	ccatggatgg	4680
416	gtgatgggg	tccctgcggc	cagtttcag	gcagagggtc	ctggcaggat	atccttctgt	4740
418	ccagtgcacc	atctggaccc	cagttccct	tcaaagggg	ggatgaccgt	gagtcctggc	4800
420	cctctgtgtt	ttataatagg	acctgcccgt	gctcaggca	cttcatgggt	ttcaactgcg	4860
422	gaaactgtaa	gtttggattt	ggggggccaa	atttacaga	gaagcgagtc	ttgatttagaa	4920
424	gaaacatttt	tgatttgagt	gtctccgaaa	agaataagtt	ctttcttac	ctcaactttag	4980
426	caaaacatac	tatcagctca	gtctatgtca	tccccacagg	cacctatggc	caaataatgaa	5040
428	atgggtcaac	accatgtttt	aatgatatac	acatctacga	cctcttggta	tggatgcatt	5100
430	actatgtgtc	aaggacaca	ctgttgggg	gctctgaaat	atggagggac	attgattttg	5160
432	cccatgaagc	accagggttt	ctgccttggc	acagactttt	cttggatttg	tgggacaaag	5220
434	aaattcggaga	actaactggg	gatgagaact	tcactgttcc	atactggat	tggagagatg	5280
436	cagaaaactg	tgacatttgc	acagatggat	acttgggagg	tcgtcaccc	gaaaatccta	5340
438	acttactcag	cccagatcc	tttttctct	cctggcagat	catttgcgt	agatcagaag	5400
440	agtataatag	ccatcagggtt	ttatgcgtat	gaacacctga	gggaccacta	ttacgtatc	5460
442	ctggaaacca	tgacaaagcc	aaaacccca	ggctccatc	ttcagcagat	gtggattttt	5520
444	gtctgagttt	gaccaggat	aatctggat	caatggatag	aactgccaat	ttcagcttta	5580
446	gaaacacact	ggaaggattt	gccagttccac	tcacaggat	agcagatcct	tctcaaagta	5640
448	gcatgcacaa	tgccttacat	atctttatga	atgaaacaat	gtcccaagta	caggatcg	5700
450	ccaaacgatcc	catttttctt	cttcaccatg	cttttgcgtt	cagtattttt	gaacaatggc	5760
452	tgcgaaggca	ccgcctctt	ttggaaagttt	accagaac	caatgcac	atcgccata	5820
454	acagagactc	ttacatggtt	ccttcatac	cgctctata	aatgggtat	ttcttcataa	5880
456	catccaagg	tctggatat	gactacagt	acctccaaga	gtcagatcca	ggcttttaca	5940
458	gaaattata	tgagcattac	ttggaaacaag	ccagtcgtat	ctggccatgg	cttcttgggg	6000
460	cagcactgg	gggagctgtt	attgtcgac	ctctctctgg	gcttagcagt	aggctatgcc	6060
462	ttcagaagaa	gaagaagaag	aagcaacccc	aggagggaaag	gcagccactc	ctcatggaca	6120
464	aagacgacta	ccacagctt	ctgtatcaga	gccatctgt	aacatcctag	gaaacagagt	6180
466	gggactgaaa	ggtttaccc	cactcgac	atttgcgtt	gtttctacaa	attaaacta	6240
468	gtataaaaca	tagaccatag	ctgttggct	tttttcaga	cccatgtttt	ttcctaagtc	6300

## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/996,128

DATE: 12/06/2001

TIME: 11:42:51

Input Set : A:\sequence listing.txt

Output Set: N:\CRF3\12062001\I996128.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No  
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date